

IMPROVED AAV VECTORS FOR GENE THERAPYBACKGROUND OF THE INVENTION

Adeno-associated virus (AAV) is a parvovirus having a single-stranded DNA genome of about 4.6 kb. Unlike other viruses, AAV is naturally defective, requiring coinfection with a helper virus (e.g. adenovirus or herpes virus) to establish a productive infection. No human disease has been found to be associated with AAV infection (Blacklow et al., 1968). The host range of AAV is broad; unlike retroviruses, AAV can infect both quiescent and dividing cells *in vitro* and *in vivo* (Flotte et al., 1993; Kaplitt et al., 1994; Podsakoff et al., 1994; Russell et al., 1994) as well as cells originating from different species and tissue types *in vitro* (Lebkowski et al., 1988; McLaughlin et al., 1988). When infection occurs in the absence of a helper virus, wild-type AAV can integrate into the cellular genome as a provirus, until it is rescued by superinfection with adenovirus. (Handa et al., 1977; Cheung et al., 1980; Laughlin et al., 1986).

The AAV genome is relatively simple, containing two open reading frames (ORFs) flanked by short inverted terminal repeats (ITRs). The ITRs contain, *inter alia*, *cis*-acting sequences required for virus replication, rescue, packaging and integration. The integration function of the ITR permits the AAV genome to integrate into a cellular chromosome after infection.

The nonstructural or replication (Rep) and the capsid (Cap) proteins are encoded by the 5' and 3' ORFs, respectively. Four related proteins are expressed from the *rep* gene; Rep78 and Rep68 are transcribed from the p5 promoter while a downstream promoter, p19, directs the expression of Rep52 and Rep40. The larger Rep proteins (Rep78/68) are directly involved in AAV replication as well as regulation of viral gene expression (for review, see

Muzyczka, 1992). The *cap* gene is transcribed from a third viral promoter, p40. The capsid is composed of three proteins of overlapping sequence; the smallest (VP-3) is the most abundant. Because the inverted terminal repeats are the only AAV sequences required in *cis* for replication, packaging, and integration (Samulski *et al.*, 1989), most AAV vectors dispense with the viral genes encoding the Rep and Cap proteins and contain only the foreign gene inserted between the terminal repeats.

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Interest in AAV as a vector in gene therapy results from several unique features of its biology. Stable genetic transformation, ideal for many of the goals of gene therapy, may be achieved by use of such AAV vectors. Furthermore, the site of integration for AAV is well-established as being on chromosome 19 of humans. This predictability removes the danger of random insertional events into the cellular genome that may activate or inactivate host genes or interrupt coding sequences, consequences that limit the use of vectors whose integration is random, e.g., retroviruses. Because the rep protein mediates the integration of AAV, it is believed that removal of this protein in the construction of AAV vectors result in altered integration patterns. In addition, AAV has not been associated with human disease, obviating many of the concerns that have been raised with virus-derived gene therapy vectors.

Notwithstanding the attractive aspects of AAV-based vectors, rapid progress in their evaluation for gene therapy has been hampered by the inability to produce recombinant viral stocks at large-scale and to high titer. The conventional method for production of recombinant AAV (rAAV) vectors is cotransfection of one plasmid containing the vector and a second helper plasmid encoding the AAV Rep and Cap proteins into 293 cells infected with adenovirus (e.g. Lebkowski *et al.*, 1988; Samulski *et al.*, 1989,

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Muzyczka, N., 1992, Kaplitt et al., 1994; Einerhand et al., 1995). This method is cumbersome and results in a low yield of rAAV, typically 10^4 - 10^5 infectious or transducing units/ml. Strategies to improve this scheme have included increasing transfection efficiency by complexing plasmid DNA to adenoviral particles via polylysine (Mamounas et al., 1995), delivering the vector sequences as part of a recombinant adenovirus (Thrasher et al., 1995) and amplification of helper plasmid copy number by linkage to a SV40 replicon (Chiorini et al. 1995).

Progress in the development of AAV as a gene therapy vector has been limited by an inability to produce high titer recombinant AAV stock using the approaches described above. The limitations to date have been thought to derive from inadequate production of the AAV proteins required in *trans* for replication and packaging of the recombinant AAV genome. *Trans*-based strategies to vector production are those that modulate the level of proteins required in *trans* to effectuate AAV vector production. Attempts to increase the levels of these proteins have included placing the AAV *rep* gene under the control of the HIV LTR promoter (Flotte, F.R. et al., *Gene Therapy* 2:29-37, 1995) to increase protein levels and the development of cell lines that express the *rep* proteins (Yang, Q. et al., *J. Virol.* 68: 4847-4856, 1994).

The limitations in producing high titer AAV vector stock may also result from a failure to include AAV *cis*-required elements in the recombinant AAV vector design. *Cis*-based strategies to increase vector production are those that provide DNA sequences required in *cis* (in tandem) with the recombinant DNA to be packaged into the AAV vector particle. The *trans* and *cis* functions are related. *Trans*-required proteins are necessary to effectuate vector production, but they require *cis*-acting sequences in the recombinant AAV genome in order to be

functionally active. Therefore, high yield AAV vector production requires a coordinated strategy of *trans*-based and *cis*-based improvements so that progress in the development of AAV as a standard gene therapy vehicle may
5 be realized.

Thus, there is a need in the art for methods and compositions which enable production of high titer recombinant AAV (rAAV) preparations that are free from
10 wild-type AAV and Adenovirus helper contamination.

SUMMARY OF THE INVENTION

15 The present invention is directed to methods for generating high titer, contaminant free, recombinant AAV vectors.

The present invention provides methods and
20 genetic constructs for producing AAV recombinant vectors conveniently and in large quantities.

The present invention further provides methods for the delivery of all essential viral proteins required
25 in *trans* for high yields of recombinant AAV.

The present invention provides recombinant AAV vectors for use in gene therapy, using *trans*- and *cis*-based strategies.
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The present invention also provides novel packaging cell lines which obviate the need for cotransfection of vector and helper plasmids.
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The invention is also directed to helper plasmids and vector plasmid backbone constructs that are used in these methods.

5 The present invention provides a reporter assay for determining AAV vector yield.

Further provided are recombinant AAV vectors in a pharmaceutically acceptable carrier.

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The present invention also provides methods of delivering a transgene of interest to a cell.

15 Compositions and methods for delivering a DNA sequence encoding a desired protein to a cell are provided by the present invention.

Still further provided are transgenic non-human mammals that express a human chromosome 19 AAV integration locus.

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DESCRIPTION OF THE FIGURES

25 FIGURE 1 shows a diagram of a replicating helper plasmid containing adenovirus genes required for AAV vector production.

FIGURE 2 shows a diagram of a replicating helper plasmid containing the AAV rep and cap genes required for AAV vector production.

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FIGURE 3 shows a diagram of a nonreplicating helper plasmid containing the AAV rep and cap genes required for AAV vector production.

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FIGURE 4 shows a diagram of a replicating helper plasmid containing the adenovirus genes and AAV rep and cap genes required for AAV vector production.

5 FIGURE 5 shows a diagram of AAV subgenomic fragments to be used in vector plasmid constructs for AAV vector production. Reference is to restriction sites in pIM45 that define the borders of the fragments.

10 FIGURE 6 shows a diagram of pTRCAT reporter plasmid.

FIGURE 7 shows a diagram of helper plasmids containing the AAV rep and cap genes used in AAV vector production.

15 FIGURE 8 shows a diagram of pTRLacZ reporter plasmid.

FIGURE 9 shows a Western blot analysis of rep protein expression from AAV nonreplicating helper plasmids. The rep proteins (in kd) are indicated at right.

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FIGURE 10 shows a Western blot analysis of cap protein expression from AAV nonreplicating helper plasmids. The cap proteins (VP, in kd) are indicated at right.

25 Figure 11 shows AAV helper plasmids represented in linear form with the thin line (only a portion of which is shown) depicting the backbone plasmid DNA, the thick bars represent the Rep and Cap coding regions and their associated control regions, the arrows above the bars depict the positions of the endogenous AAV promoters, p5, p19 and p40, and the "X" indicates that the p40 promoter has been inactivated by mutation.

30 Figure 12 shows Rep and Cap gene expression from AAV helper plasmids analyzed by Western blot using anti-Rep (panel A) and anti-Cap (panel B) primary antibodies. Lanes in each panel correspond to samples derived from cells transfected

WO 97/09441

with the following helper DNAs: lane 1 (mock), lane 2 (pIM45), lane 3 (pIMRSV), lane 4 (p5repΔ-CMVcap), lane 5 (pRSV repΔ-CMVcap), lane 6 (p5repΔ-p40cap), lane 7 (pRSVrepΔ-p40cap), lane 8 (pIMRSV-am), lane 9 (wtAAV) MOI=10. Molecular weight size standards (in kD) are depicted to the left of each panel; each of the AAV Rep and Cap proteins is identified to the right.

Figure 13 is an analysis of the mechanism of down-regulation of the cap genes. Shown is a Western blot analysis of samples derived from 293 cells transfected with the appropriate DNAs (10 µg total) in the absence (-) or presence (+) of adenovirus (AD5ts149, MOI=20). Panel A and panel B show blots developed using an anti-Rep and anti-Cap primary antibodies, respectively. Lanes in each panel correspond to the following: lane 1 (mock transfected cells), lane 2, (pIM45), lane 3 (pIMRSV), lane 4 (pIMRSV-am), lane 5 (pIMRSV-am and suppressor tRNA plasmid), lane 6 (pIMRSV-am and pRSVRep), lane 7 (pRSVRep alone; *Wt*= cells infected with wtAAV (MOI=15) in the presence of adenovirus (Adts149, MOI=20). Molecular weight size markers (in kD) are shown at the left and the AAV Rep and Cap proteins are identified at the right.

Figure 14 is an analysis of total RNA derived from transfections described in Figure 13. Panel A shows the Northern, panel B is the ethidium stained gel to demonstrate that equal amounts of RNA were loaded in each lane. Lanes in each panel correspond to the following: lane 1 (mock transfected cells), lane 2, (pIM45), lane 3 (pIMRSV), lane 4 (pIMRSV-am), lane 5 (pIMRSV-am and suppressor tRNA plasmid), lane 6 (pIMRSV-am and pRSVRep), lane 7 (pRSVRep alone; *Wt*= cells infected with wtAAV (MOI=15) in the presence of adenovirus (Adts149, MOI=20). Transfections were carried out in the absence (-) or presence (+) of adenovirus (AD5ts149, MOI=20). RNA size standards (in kilobases) are shown at the left in panel A.

at the right in panel B; AAV mRNAs are identified at the right of panel A.

Figure 15 is an analysis of vector replication and levels of wt AAV contamination. Adenovirus (Ad5ts149) - infected 293 cells were transfected with 1.5 µg of vector (pTRLacZ) and 15 µg of the indicated helper DNA. The replicated viral DNA was then analyzed by Southern blot; duplicate filters were probed with the lacZ probe. Panels C and D show primary and secondary Hirt DNAs, respectively, probed with the AAV fragment. For the primary DNAs (panels A and C), lanes correspond to the following samples: lane 1 (mock transfection), lane 2 (pIM45), lane 3 (pIMRSV), lane 4 (p5repΔ-CMVcap), lane 5 (RSVrepΔ-CMVcap), lane 6 (p5repΔ-p40cap), lane 7 (RSVrepΔ-p40cap), lane 8 (pIMRSV-am). For the secondary DNAs (panels B and D), the lanes are the same except the sample in lane 8 is derived from cells infected with wtAAV (MOI=0.001) and adenovirus (Ad5ts149, MOI=20). The positions of DNA size standards in (kilobase pairs) are depicted at the left of each panel; hybridizing bands corresponding to the dimer replicative form (dRF) and monomer replicative form (mRF) are identified at the right.

DETAILED DESCRIPTION OF THE INVENTION

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All patent applications, patents, and literature references cited in this specification are hereby incorporated by reference in their entirety. In case of conflict or inconsistency, the present description, including definitions, will control.

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Definitions:

293 cells - human embryonic kidney cell line harboring and expressing parts of the adenovirus genome including the adenoviral E1 region.

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293-MT-DBP cells - human embryonic kidney cell line modified to express parts of the adenovirus genome which complement recombinant adenovirus vectors that are deleted for E1 and E2A. Deposited August 28, 1996 with the American Type Culture Collection, 12301 Parklawn Drive, Rockville, Maryland, and assigned ATCC CRL -12181.

2C4 cells - human embryonic kidney cell line modified to express parts of the adenovirus genome which complement recombinant adenovirus vectors that are deleted for E1 and E4. Deposited August 28, 1996 with the American Type Culture Collection, 12301 Parklawn Drive, Rockville, Maryland, and assigned ATCC CRL -12182.

3B1 cells - human embryonic kidney cell line modified to express parts of the adenovirus genome which complement recombinant adenovirus vectors that are deleted for E1 and E2A. Deposited August 28, 1996 with the American Type Culture Collection, 12301 Parklawn Drive, Rockville, Maryland, and assigned ATCC CRL -12183.

Cfu - colony forming units. For retroviral and adeno-associated virus vectors carrying antibiotic resistance genes, the number of antibiotic-resistant cell colonies after infection. It is assumed that one colony arises from a single infected cell.

Expression plasmid - extrachromosomal genetic elements that can propagate autonomously in cells, constructed in such a way that the genes carried in the plasmid can be expressed in the cells.

In cis - from the same DNA molecule.

In trans - from a different DNA molecule.

Insertional mutagenesis - the introduction of a mutation into a target gene by the insertion of foreign DNA, such as viral DNA, into the host-cell genome. The development of a mutation in a cellular gene as the result of the introduction of foreign DNA into a cell.

ITR - inverted terminal repeat, a DNA sequence that is repeated at either end of the virus in an inverted form.

Promoter-transgene cassette - a combination of DNA sequences containing elements necessary for directing production of a gene product and the DNA sequence of the gene itself.

Transduction - the introduction of foreign DNA into cells of an organism (*in vivo*).

Transfection - the introduction of foreign DNA into cells in culture (*in vitro*). Genetic modification of eukaryotic cells by introduction of foreign DNA using chemical means. In transient transfection, expression occurs from unintegrated foreign DNA and can be detected for a few days after transfection.

Transgene - a gene that has been stably incorporated into another organism.

Quiescent cells - cells that are not actively dividing.

Recombination - physical interaction between two or more DNA molecules, in this case viral sequences, leading to exchange of DNA sequence between the molecules.

Titer - the number of virus particles produced per ml. The assay system to determine the number of virus

particles produced varies considerably depending on the virus in question. High titers are generally essential for successful gene therapy since they allow introduction of the therapeutic gene carried by the virus into the maximum number of cells.

Vector - a vehicle, usually a biological entity, such as a virus, used for the delivery of genes into an organism. A reagent that facilitates the introduction of foreign genes into cells.

LacZ gene - bacterial gene used as a cellular gene marker. Its expression is detected by a blue coloration in the presence of the substrate X-gal.

Packaging cells - cells that have been transfected with plasmids containing the cap and rep genes from AAV.

A purpose of the investigation described herein was to determine the limiting component(s) required for rAAV packaging. An understanding of the process at a basic level should benefit all methods of rAAV production. By selectively increasing expression of either the rep or cap genes (or both), it is shown that Cap protein production is one limiting factor in the production of rAAV.

A. PROVISION OF TRANS PRODUCT

30 1. Adenovirus Proteins

As discussed, adenovirus proteins are required to generate a productive AAV infection. In the absence of helper adenovirus, AAV integrates into the cellular genome, remaining latent until the cell is infected with adenovirus.

Adenovirus genes required for helper function include, *inter alia*, E1A, E1B, E2A, E4 ORF6, and VA RNA (Muzycka, N., *Curr. Top. Micro. Immunol.* 158: 97-129, 1992). Standard methods of generating recombinant AAV
5 vectors have relied on adenovirus infection of the target cell in order to provide adequate levels of necessary helper proteins. To avoid the unwanted generation of wild-type adenovirus that may occur during AAV vector
10 production, an adenovirus containing a deletion is used in a cell line that provides an essential adenovirus protein *in trans*. Alternatively, a temperature-sensitive adenovirus replication mutant may be used at a nonpermissive temperature.

15 In one embodiment of the present invention, a helper plasmid is provided which contains the essential adenovirus helper genes bounded by AAV ITR sequences that allow the plasmid to replicate. The helper plasmid may contain E1A, E1B, E2A, E4 ORF6, and VA RNA genes. Each of
20 these genes may also have its own promoter, through which transcription can occur. Alternative promoters that may be used in the helper plasmid include, but are not limited to, CMV, RSV, MMTV, E1A, EF1a, actin, cytokeratin 14, cytokeratin 18, PGK as well as others known to those
25 skilled in the art.

Providing the essential adenovirus genes on a replicating plasmid is an alternative to infecting a host cell with adenovirus. The AAV ITR sequences in the plasmid
30 function as an origin of replication under the control of the AAV rep proteins to increase the plasmid copy number. Increased copy number leads to increased levels of the proteins encoded by the genes on the plasmid. Thus, the helper plasmid of the present invention provides the
35 adenovirus proteins required for AAV vector production, while eliminating the possibility of adenovirus production. A further advantage is that the levels of the adenovirus

proteins are not limited by the amount of input plasmid DNA, since replication of the plasmid will increase gene copy number above input levels.

5 In additional embodiments, the origin of replication may include, but is not limited to, the SV40 origin of replication, the Epstein-Barr (EBV) origin of replication, as well as others known to those skilled in the art. Where, for example, an origin of replication
10 requires an activating protein - e.g., SV40 origin requiring T antigen, EBV origin requiring EBNA protein - the activating protein may be provided by stable transfection so as to create a cell line source, or by transient transfection with a plasmid containing the
15 appropriate gene.

Standard recombinant DNA techniques may be employed to construct the helper plasmids of the present invention (see, e.g., *Current Protocols in Molecular Biology*, Ausubel., F. et al., eds, Wiley and Sons, New York
20 1995). Such methods include the utilization of compatible restriction sites at the borders of the adenovirus genes and the ITR sequences or DNA linker sequences which contain restriction sites, as well as other methods known to those
25 skilled in the art. Reference for adenovirus DNA sequence information is given in Roberts, R.J., in *Adenovirus DNA: The Viral Genome and Its Expression*, Oberfler, W., ed., Martinus Nihoff Publishing, Boston, 1986). Plasmids routinely employed in molecular biology - e.g., pBR322 (New
30 England Biolabs, Beverly, MA), pRep9 (Invitrogen, San Diego, CA) or pBS (Stratagene, La Jolla, CA) - may be used as the basis for the helper plasmid into which adenovirus genes and the AAV ITR may be inserted. The adenovirus genes may be placed into the helper plasmid in any
35 positional order. A particular embodiment of such a replicating helper plasmid according to the invention is shown in FIGURE 1.

The helper plasmid may be used in the generation of recombinant AAV when combined with a source of the AAV rep and cap proteins, as well as the recombinant AAV genome. Transfection of cells by the plasmid using techniques well known in the art will provide the adenovirus gene products necessary for initiation of AAV rep gene expression.

10 2. AAV Proteins

In order to generate recombinant AAV vector stocks, standard approaches provide the AAV rep and cap gene products on a plasmid that is used to cotransfect a target cell along with the AAV vector plasmid. The levels of rep and cap proteins produced as a result of transfection are relevant to maximizing AAV vector production. This is because the rep proteins activate transcription of the cap gene, leading to production of the AAV structural proteins that are involved in packaging the recombinant genome.

Attempts to increase the levels of these AAV proteins in order to enhance vector production have been problematic (Kotin, R.M., *Human Gene Therapy* 5:793-81, 1994). One of the problems appears to be the toxicity of the rep protein to the cell at high levels.

In this aspect of the present invention, AAV rep and cap genes are provided on a replicating helper plasmid that contains the AAV ITR sequences. The rep proteins activate ITR as an origin of replication, leading to replication of the plasmid, that result in increased copy number. The advantage of this method is that rep protein level is not simply dependent on the efficiency of transfection with the plasmid, but is also a function of the replication of the plasmid following transfection. An

example of a replicating helper plasmid containing the AAV rep and cap genes is provided in FIGURE 2. In other embodiments of this aspect of the invention, the origin of replication may include, but is not limited to, the SV40 origin of replication, the Epstein-Barr (EBV) origin of replication, as well as others known to those skilled in the art. Where, for example, an origin of replication requires an activating protein - e.g., SV40 origin requiring T antigen, EBV origin requiring EBNA protein - the activating protein may be provided by stable transfection so as to create a cell line source, or by transient transfection with a plasmid containing the appropriate gene.

In still another embodiment of the invention, AAV rep and cap genes may be provided on a non-replicating plasmid, which does not contain an origin of replication. Such non-replicating plasmid further insures that the replication apparatus of the cell is directed to replicating recombinant AAV genomes, in order to optimize production of virus. Furthermore, since some studies have suggested that high levels of rep protein may be toxic to the cell (Muzyczka, N., *Curr. Top. Micro. Immunol.* 158: 97-129, 1992), providing the rep gene on a non-replicating plasmid may decrease this possibility. The levels of the AAV proteins encoding by such non-replicating plasmids may be modulated by use of particular promoters to drive the expression of these genes. Such promoters include, *inter alia*, AAV promoters, as well as promoters from exogenous sources, e.g., CMV, RSV, MMTV, E1A, EF1a, actin, cytokeratin 14, cytokeratin 18, PGK, as well as others known to those skilled in the art. An example of a non-replicating AAV helper plasmid is provided in FIGURE 3.

Levels of rep and cap proteins produced by these helper plasmids may be individually regulated by the choice of a promoter for each gene that is optimally suited to the

level of protein desired. Specific modulation of a particular gene - e.g., the rep gene - may also be achieved with the use of an inducible promoter. Such inducible promoters include, but are not limited to, MMTV, 5 metallothionein, as well as others known to those skilled in the art.

In order to prevent any packaging of AAV genomic sequences containing the rep and cap genes, a plasmid 10 containing the rep and cap DNA fragment may be modified by the inclusion of a "stuffer" fragment into the AAV genome which causes the DNA to exceed the length for optimal packaging. Thus, the helper fragment is not packaged into AAV virions. This is a safety feature, ensuring that only 15 a recombinant AAV vector genome that does not exceed optimal packaging size is packaged into virions. An AAV helper fragment that incorporates a stuffer sequence exceeds the wild-type genome length of 4.6 kb, and lengths above 105% of the wild-type will not be packaged. The 20 stuffer fragment may be derived from, for example, such non-viral sources as the β -galactosidase gene.

Standard recombinant DNA techniques may be employed to construct the helper plasmids of the present 25 invention (see e.g., *Current Protocols in Molecular Biology*, Ausubel., F. et al., eds, Wiley and Sons, New York 1995), including the utilization of compatible restriction sites at the borders of the genes and AAV ITR sequences (where used) or DNA linker sequences which contain 30 restriction sites, as well as other methods known to those skilled in the art. Reference for AAV DNA sequence information is given in Srivastava, A. et al., *J. Virol.* 45:555-564, 1983. Plasmids routinely employed in molecular biology may be used as a backbone - e.g., pBR322 (New 35 England Biolabs, Beverly, MA), pRep9 (Invitrogen, San Diego, CA), pBS (Stratagene, La Jolla, CA) - for the

insertion of the AAV genes and, in the case of a replicating plasmid, the AAV ITR.

3. Hybrid Helper Plasmid

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Generation of a recombinant AAV vector stock requires both AAV and adenovirus proteins, provided in trans, in order to facilitate transcriptional activation, replication and packaging of the recombinant AAV genome. Standard approaches have utilized plasmid-based delivery of AAV genes into the target cell. Infection of the target cell with adenovirus is used to provide adenovirus genes. This multi-step protocol requires coordination of transfection and infection. Furthermore, infection of the cell with adenovirus allows for adenovirus production, which is not desirable when attempting to produce a pure AAV vector stock. In addition, the introduction of many viral genes which are not needed in vector generation causes diversion of transcriptional and replication machinery that could be directed to more efficient production of the essential proteins for AAV production. While AAV vectors have been produced using adenovirus genes introduced by infection, high yield vector production remains problematic. Therefore, a more efficient delivery of the genes coding for the proteins required in trans should improve AAV vector production.

The present invention provides a simple method for the delivery of all essential viral proteins required in trans for high yields of AAV. A hybrid plasmid is constructed to carry AAV and adenovirus genes encoding the essential proteins. The advantages of such a plasmid include, *inter alia*, a single entry event into the cell delivering all genes coding for trans-acting proteins, coordinate provision of all such genes and avoidance of adenovirus production resulting from the elimination of unnecessary adenovirus genes. Such a plasmid is shown in

FIGURE 4. The plasmid contains essential adenovirus genes - E1A, E1B, E2A, E4 ORF6, and VA RNA. The plasmid also contains the AAV *rep* and *cap* genes, as well as the AAV ITR sequences which are required to replicate the plasmid. The genes may be transcribed using their own promoters. Alternatively, promoters may include, but are not limited to, CMV, RSV, MMTV, E1A, EF1a, actin, cytokeratin 14, cytokeratin 18, PGK, as well as others known to those skilled in the art. The adenovirus genes may be inserted into the plasmid in the order shown in FIGURE 4, or they may be inserted in any number of different positional arrangements that are within the ability of the skilled artisan to devise.

All essential genes required in *trans* may also be provided on two plasmids for ease of cloning. Thus, the AAV and adenovirus genes on the hybrid helper plasmid may be carried by two plasmids, in any optimal arrangement for cloning. In other words, the AAV and adenovirus genes do not have to be on separate plasmids.

Standard recombinant DNA techniques may be employed to construct the helper plasmids of the present invention (see, e.g., *Current Protocols in Molecular Biology*, Ausubel., F. et al., eds, Wiley and Sons, New York 1995), including the utilization of compatible restriction sites at the borders of the genes and the ITR sequences or DNA linker sequences which contain restriction sites, as well as other methods known to those skilled in the art. Reference for adenovirus and AAV DNA sequence information is cited above. Routinely used plasmids - e.g., pBR322 (New England Biolabs, Beverly, MA), pRep9 (Invitrogen, San Diego, CA), pBS (Stratagene, La Jolla, CA) - may be used for the insertion of the adenovirus and AAV genes and the AAV ITR. The adenovirus genes may be placed into the helper plasmid in any positional order.

4. Production of Recombinant AAV Vectors

Helper plasmids that provide essential proteins required in trans are used to generate recombinant AAV vector stock. These plasmids are introduced into the target cell using any number of transfection methods, including, *inter alia*, calcium-phosphate transfection, lipofection or other techniques known to those skilled in the art. The ratio of helper plasmids to the quantity of vector plasmid containing the gene of interest range from 1:1-1:10. This procedure produces recombinant AAV vectors; the vector plasmid contains the recombinant AAV genome flanked by the AAV ITRs.

Recombinant AAV vectors are produced using 293 cells in 8 roller bottles (1×10^9 cells/ml). Cells are transfected with both the helper plasmid and the AAV vector plasmid at a vector:helper ratio of 1:10. The plasmids may be introduced into the target cell using any number of transfection methods, including, but not limited to, calcium-phosphate, lipofection, or other techniques known to those skilled in the art (see e.g., *Current Protocols in Molecular Biology*, Ausubel, F. et al., eds., Wiley and Sons, New York, 1995). In a preferred embodiment, lipofection is used for transfection. Adenovirus infection is initiated at a multiplicity of infection (MOI) of 20. The adenovirus strain may be a deletion virus, in which case, complementing genes are integrated into the cell line, or a temperature-sensitive mutant (e.g., ts149) which cannot replicate at a nonpermissive temperature (39°C). The transfected/infected cells are then incubated for 2 days at the appropriate temperature. After incubation, the cells are harvested and then lysed by three (3) freezing and thawing cycles in the presence of benzonase (American International Chemical, Natick, MA). 1% deoxycholate and 0.25% trypsin is then added to the

lysate, followed by incubation for 30 minutes at 37°C. The cell lysate (2 roller bottles/gradient) may then be applied to a CsCl step gradient (1.5g/ml-1.36g/ml) in a SW28 rotor and centrifuged at 26K for 6 hours at 4°C. Fractions are obtained and then further purified on two equilibrium gradients, using a NVT65 rotor, and centrifuged at 60K for 20 hours at 4°C. Fractions from the equilibrium gradients are screened on a refractometer and pooled. Pooled fractions are dialyzed in PBS with 1% sucrose 3 times for 2 hours at 4°C.

5. Assay for Recombinant AAV Vector Production

The efficiency of a helper plasmid as a source of proteins required in trans is determined from the yield of AAV vector stock. To determine virus yield, an AAV infectious center assay is used to assay for production of infectious progeny. The recombinant AAV vectors are recovered after production using the purification protocol described above. The assay shows whether infectious AAV progeny are produced in the production protocol.

The assay is performed with 293 cells that are plated on day one at a density of 1×10^5 cells per well in 0.1 ml/well in DME medium supplemented with 10% FBS, penicillin/streptomycin, and glutamine. After about 24 hours, the cells are infected with adenovirus at an MOI of 20 and with wild-type AAV at an MOI of 2. The viruses are suspended in the same medium, and 0.1 ml is added to each well to initiate infection. The AAV vector samples are added to the well (25-100 microliters of lysate or dilutions) and the plates are incubated for 24-30 hours at the appropriate temperature (37°C for wild-type adenovirus; 39°C for an adenovirus temperature-sensitive mutant). On the day after infection, the medium is carefully removed from the cells. Cold 0.2 ml PBS containing 5mM EDTA is added to each well, and the plate is put on ice.

5 by placing a nitrocellulose filter prewetted with PBS in position, and adding 5 ml of PBS to the top of the filtration unit. The cells in the plate are resuspended by pipetting. 0.05 ml of the cell suspension is added into the PBS buffer in the filtration unit and mixed by rotation. Suction is applied to the apparatus to deposit the cells onto the filters. The filters are air dried for 10 minutes. The cells are lysed on directly on the filters using denaturing solution followed by neutralizing solution. The filters are dried on paper for 5 minutes following each solution and then air dried for 10 minutes. The filters are washed in 2X SSC, air dried for 10 minutes, and baked in a vacuum oven for 2 hours. Hybridization to a probe that detects the gene of interest is performed using the filters prepared as above, wetted in 2X SSC. Filters can be prehybridized using 30-40 ml of Quick-Hyb® (Stratagene, La Jolla, CA) by incubating at 68°C for 2-4 hours in a rotating water bath. The labelled probe is then added to the Quick-Hyb® solution, and incubated overnight at 68°C. Filters are washed the next day (5 minutes in 2X SSC, 0.5% SDS at room temperature, followed by 2 hours in 0.1X SSC, 0.5% SDS at 65°C). The filter is exposed to film overnight at -80°C to produce an autoradiograph.

25 The number of infectious centers on the filter are counted on the autoradiograph. The titer of the starting material is determined by multiplying the number of infectious centers by any dilution used in the preparation of test samples.

30 Where the AAV vector in production contains a reporter gene, alternative methods for determining the AAV vector titer can be used. For example, if the lacZ gene is used, the infected cells can be stained for the expression

of the gene product, β -galactosidase, using X-gal. Titer is determined, for example, by counting the blue-stained cells in a plate well.

5 B. AAV VECTOR PLASMID CONSTRUCTS CONTAINING CIS-ACTING SEQUENCES

 The present invention also provides a means to increase the production of recombinant AAV vectors via AAV
10 vector plasmid designs that utilize *cis*-acting sequences in the AAV genome required for efficient replication and packaging. The invention also provides vector plasmids designed to provide such *cis*-acting sequences.

15 Current vector plasmid designs place the gene of interest between the AAV ITR sequences to create a recombinant genome and provide no other AAV sequences. The AAV ITR sequences have *cis*-acting functions that facilitate replication and packaging of the recombinant genome during
20 vector production, as well as integration of the vector DNA into a cell following its introduction by the AAV vector. Thus, ITR sequences are retained in recombinant AAV vector designs. However, the difficulty in achieving high titer production of AAV vectors indicates that the ITRs *per se*
25 are not sufficient to provide all *cis*-acting functions necessary to the production of high titer vector stock. Therefore, other *cis*-acting AAV sequences in addition to the ITRs are required within the vector construct in order to increase the efficiency of replication and/or packaging
30 of the recombinant AAV genome.

Cis-acting elements in the AAV genome are believed to facilitate rescue and replication of the genome through interactions with the AAV rep proteins. It is
35 known that rep proteins bind to sites in the AAV ITR as well as to sites in the AAV p5 and p19 promoters (McCarty, D.M. et al., *J. Virol.* 65: 2936-2945, 1991; McCarty, D.M.

et al., *J. Virol.* 68:4988-4997, 1995). *Cis*-acting packaging elements also appear to be required in the recombinant AAV vector genome for maximal particle production.

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The present invention provides a method to improve AAV vector production using vector backbone constructs that contain AAV sequences in addition to the ITR sequences. The AAV vector backbone may include AAV genomic fragments that contain *rep*-binding sites or critical packaging sequences. Because the precise number and location of all *cis*-acting AAV sequences has not yet been defined, construction of vector plasmids containing significant portions of the AAV genome is important to include all *cis*-acting sequences, including those that are still undefined. While these vector plasmid constructs improve the production of recombinant AAV vector stock, a further utility of the invention is that essential *cis*-acting sequences can be functionally identified through improved vector production.

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The vector constructs containing such *cis*-acting sequences may be prepared using known techniques. (see e.g. *Current Protocols in Molecular Biology*, Ausubel., F. et al., eds, Wiley and Sons, New York 1995). The presence of known restriction sites in the AAV genome may be used to derive subgenomic fragments for insertion into a recombinant AAV vector. Fragment length is chosen so that the recombinant genome does not exceed the packaging capacity of the AAV particle. If necessary, a "stuffer" DNA sequence is added to the construct to maintain standard AAV genome size for comparative purposes. Such a fragment may be derived from such non-viral sources, e.g., *lacZ*, or other genes which are known and available to those skilled in the art.

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The present invention provides a series of vector plasmid constructs which add AAV subgenomic fragments into a vector plasmid containing a gene of interest flanked by the AAV ITRs. See FIGURE 5. These fragments range in size
5 from 1.7 - 2.1 kb. Because these fragments contain coding as well as noncoding regions of the AAV genome, effects on packaging may be due to elements acting in trans as well as in cis. Further modifications of the constructs shown -
10 e.g., insertion of smaller AAV fragments - are within the scope of the invention, and may be readily made by those skilled in the art. When subgenomic fragments are used that are smaller than the coding regions in the genome, the effects observed on vector production are characterized as cis-acting effects. For example, standard deletion
15 analysis approaches are suitable to refine subgenomic fragments to a minimal length needed for optimal vector production. Defined AAV cis-acting fragments - e.g., responsive elements - are specifically cloned into the vector plasmids.

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The present invention provides an efficient reporter assay for determining AAV vector yield for use in gene therapy. In this manner, the most efficient construct designs are identified by production of high titer stock.

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A plasmid containing a reporter gene and the AAV ITR sequences is used to determine improvement in production efficiency when AAV sequences are added. This plasmid may be modified with the insertion of AAV
30 subgenomic fragments to create additional constructs, such as plasmid pTR-CAT, shown in FIGURE 6. This plasmid contains an expression cassette comprising the chloramphenicol acetyltransferase (CAT) gene under the control of CMV promoter, a hybrid intron and the BGH polyA
35 site. The expression cassette was cloned into pUC-TR, a high copy number plasmid derived by cloning the AAV ITR region into pUC19. (New England Biolabs, Beverly, MA) The

insertion of the expression cassette into pUC-TR created pTR-CAT. Restriction sites may be added at the 5' and 3' ends of the CAT cassette by inserting linkers or by using other techniques known to those skilled in the art. AAV subgenomic fragments may be inserted at these restriction sites. This reporter plasmid will accommodate the subgenomic fragments shown in FIGURE 5 that may be placed at either the 5' or 3' end of the expression cassette. Optimally, it may be necessary to test the insertion of fragments at both ends in order to determine any positional effect that the placement of the insert might have. The fragment can also be inserted at both ends of the vector plasmid. One or more copies of a defined *cis*-acting element can increase the amount of vector produced.

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The packaging of the novel constructs requires that of AAV *rep* and *cap* genes (e.g., using pIM45, described in EXAMPLE 1) be provided and as well as essential adenovirus genes. A plasmid containing the AAV genes is cotransfected with the novel construct. An adenovirus infection or an adenovirus helper plasmid of the present invention (See Section 1, *supra*) provides the other necessary genes.

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An infectious center assay utilizing an appropriate probe is used for determining the amount of infectious progeny (see Section 5, *supra*). Alternatively, a reporter gene product in the AAV vector can be assayed directly - e.g., a CAT enzyme assay is used where this reporter gene is present, for example, pTR-CAT (*Current Protocols in Molecular Biology*, Ausubel, F. et al., eds., Wiley and Sons, New York, 1995).

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Alternative reporter genes may be used in assaying the vector plasmid constructs provided that the final plasmid construct containing the reporter gene and the AAV *cis*-acting fragments do not exceed the packaging

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length for an AAV particle. Other reporter gene products may be detected in the infectious AAV particles using appropriate biochemical assay.

- 5 The invention provides transgenic non-human mammals capable of expressing the AAV integration locus on human chromosome 19. Examples of non-human transgenic mammals are transgenic cows, sheep, goats, pigs, rabbits, rats and mice.
- 10 Animal model systems which elucidate the physiological and behavioral roles of invention polypeptides are produced by creating transgenic animals in which the expression of a polypeptide of interest is
- 15 altered or modified using a variety of techniques. Examples of such techniques include the insertion of normal or mutant versions of nucleic acids encoding a polypeptide of interest, by microinjection, retroviral infection or other means well known to those skilled in the art, into
- 20 appropriate fertilized embryos to produce a transgenic animal. See, for example, Carver, et al., *Bio/Technology* 11:1263-1270, 1993; Carver et al., *Cytotechnology* 9:77-84, 1992; Clark et al., *Bio/Technology* 7:487-492, 1989; Simons et al., *Bio/Technology* 6:179-183, 1988; Swanson et al.,
- 25 *Bio/Technology* 10:557-559, 1992; Velander et al., *Proc. Natl. Acad. Sci. USA* 89:12003-12007, 1992; Hammer et al., *Nature* 315:680-683, 1985; Krimpenfort et al., *Bio/Technology* 9:844-847, 1991; Ebert et al., *Bio/Technology* 9:835-838, 1991; Simons et al., *Nature*
- 30 328:530-532, 1987; Pittius et al., *Proc. Natl. Acad. Sci. USA* 85:5874-5878, 1988; Greenberg et al., *Proc. Natl. Acad. Sci. USA* 88:8327-8331, 1991; Whitelaw et al., *Transg. Res.* 1:3-13, 1991; Gordon et al., *Bio/Technology* 5:1183-1187, 1987; Grosveld et al., *Cell* 51:975-985, 1987; Brinster et
- 35 al., *Proc. Natl. Acad. Sci. USA* 88:478-482, 1991; Brinster et al., *Proc. Natl. Acad. Sci. USA* 85:836-840, 1988; Brinster et al., *Proc. Natl. Acad. Sci. USA* 82:4438-4442,

1985; Al-Shawi et al., *Mol. Cell. Biol.* 10(3):1192-1198,
1990; Van Der Putten et al., *Proc. Natl. Acad. Sci. USA*
82:6148-6152, 1985; Thompson et al., *Cell* 56:313-321, 1989;
Gordon et al., *Science* 214:1244-1246, 1981; and Hogan et
5 al., *Manipulating the Mouse Embryo: A Laboratory Manual*
(Cold Spring Harbor Laboratory, 1986).

Another technique, homologous recombination of
mutant or normal versions of these genes with the native
10 gene locus in transgenic animals, may be used to alter the
regulation of expression or the structure of the
polypeptide of interest (see, Capecchi et al., *Science*
244:1288, 1989; Zimmer et al., *Nature* 338:150, 1989).
Homologous recombination techniques are well known in the
15 art. Homologous recombination replaces the native
(endogenous) gene with a recombinant or mutated gene to
produce an animal that cannot express native (endogenous)
protein but can express, for example, a recombinant protein
which results in expression, for example, of the human AAV
20 integration locus.

In contrast to homologous recombination,
microinjection adds genes to the host genome, without
removing host genes. Microinjection can produce a
25 transgenic animal that is capable of expressing both
endogenous and exogenous polypeptides. Inducible promoters
can be linked to the coding region of the nucleic acids to
provide a means to regulate expression of the transgene.
Tissue-specific regulatory elements can be linked to the
30 coding region to permit tissue-specific expression of the
transgene. Transgenic animal model systems are useful for
in vivo screening of vector compositions for identification
and confirmation of integration and long-term transgene
expression.

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C. CONSTRUCTION OF AAV HELPER PLASMIDS.

A series of helper plasmids was constructed to determine if rAAV packaging is limited by expression levels of the *rep* and/or *cap* genes (Figure 11). Expression of Rep and Cap proteins was increased by replacing the endogenous AAV promoters, p5 and p40, with the RSV LTR and the CMV IE promoter, respectively. The starting helper plasmid, pIM45 (McCarty et al., 1991), contains a sequence encompassing the wild-type AAV genome but excluding the terminal repeats (nucleotides 145-4493). pIMRSV is a modification of pIM45 in which the RSV LTR replaces p5. Because p40 is located within the Rep coding region, the *rep* and *cap* genes were separated to allow replacement of p40 with the CMV IE promoter (as in p5repΔ-CMVcap). This strategy generated a vector with a direct repeat of 431 bp of sequence downstream from the p40 and CMV promoter. To prevent generation of wild type AAV through recombination, the p40 promoter lying within the *rep* ORF of this construct was inactivated by site-directed mutagenesis. p5repΔ-p40cap was constructed to express the *rep* and *cap* genes from endogenous AAV promoters as in pIM45, but so as to be more directly comparable to p5repΔ-CMVcap, the Rep and Cap coding regions were separated. RSVrepΔ-CMVcap and RSVrepΔ-p40cap are derivatives of p5repΔ-CMVcap and p5repΔ-p40cap, respectively in which p5 is replaced by the RSV LTR.

D. REP AND CAP GENE EXPRESSION FROM AAV HELPER PLASMIDS.

The amounts of Rep and Cap proteins expressed from each of the AAV helper plasmids were estimated by Western blot analysis (Figure 12). The four Rep proteins produced following transfection into 293 cells in the presence of an Adts149 infection (MOI = 20) comigrate with the corresponding proteins detected after coinfection of 293 cells with wild-type AAV (wtAAV) and Adts149. For each of the helper plasmids, Rep78 and Rep52 are the major

proteins produced. Rep68 and Rep40, which are translated from spliced messages, were observed at a lower level. These were also detected as minor proteins in the wtAAV infection.

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When the p5 promoter was replaced with the RSV LTR, an increase in the level of Rep78 was observed. This was the case for all three helpers, pIMRSV, RSVrepΔ-CMVcap and RSVrepΔ-p40cap. There was no change in the amount of Rep52, because in all constructs it was derived from a p19 transcript.

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The three capsid proteins VP1, VP2, and VP3 were produced from all helper plasmids in the 1:1:10 ratio observed in a wt AAV infection (Figure 12). Synthesis of all three capsid proteins was enhanced when p40 was replaced by the CMV IE promoter (Figure 12 lane 2 vs. lane 4). However, expression of the *rep* gene from the RSV LTR appeared to have a down-regulatory effect on *cap* expression from p40. Thus, the level of capsid protein was reduced for pIMRSV relative to the parental plasmid containing p5 as the promoter regulating expression of *rep* (pIM45; compare lanes 3 and 2). A similar, but less dramatic effect was observed with capsid protein expression from the CMV IE promoter (lane 5 vs lane 4). In the latter case, a corresponding reduction was also seen in *cap* mRNA by Northern analysis, suggesting that overexpression of Rep78 results in transcriptional down-regulation of the CMV promoter. Cap protein synthesis was also reduced relative to pIM45 when the *rep* and *cap* genes were expressed from separate transcription units as in p5repΔ-p40cap where the AAVpA signal separates the *rep* and *cap* ORFs.

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It should be noted that the total level of AAV proteins produced in the transient transfections was comparable to that observed in a wt AAV infection at an MOI of 10. While Rep78, Rep52 and the capsid proteins appeared

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at levels similar to that observed in the wt AAV infection when expressed from the AAV promoters (p5, p19 and p40, respectively), expression from the heterologous promoters, RSV LTR and the CMV IE, increased the amount above that observed in the viral infection. This is especially significant when it is considered that transfection efficiency ranges from 20-50% while infection at an MOI of 10 should occur with greater efficiency. This suggests that the concentration of each viral gene product per transfected cell is higher in the transient transfections than in the wt AAV infection.

E. CONSTRUCTION OF PIM-RSVAM AND ANALYSIS OF CAP GENE DOWN-REGULATION.

To analyze further the mechanism of down-regulation of capsid protein expression in the helper plasmids containing an RSV LTR-*rep* gene cassette, a derivative of pIMRSV was made that contained an amber mutation within the *rep* ORF (pIMRSV-*am*). If down-regulation were due to an alteration in *cis* (i.e. replacement of p5 with the RSV LTR), then it should persist in the amber mutant. In contrast, the down-regulatory effect should be relieved in the mutant if it is dependent upon synthesis of full-length Rep protein.

Transient transfections in 293 cells were performed using pIMRSV-*am* as a helper in the presence and absence of adenovirus (Ad) infection. Nuclear proteins were isolated and analyzed by Western blot (Figure 13). With pIM45 as helper Rep78 appears at a high level in the absence of Ad due to expression of the E1A and E1B genes in 293 cells, hence the apparent lack of induction with the addition of Ad (induction of Rep78 with this construct is observed in HeLa cells). Infection with Ad does result in the appearance of the spliced Rep proteins, Rep68 and Rep40, and a slight increase in the level of Rep52. As

expected, expression of Rep78 from the RSV promoter is unresponsive to Ad infection; the same high level appears in the presence and absence of helper virus coinfection. In cells transfected with pIMRSV-*am*, a small amount of
5 full-length Rep78 is observed, indicating that the mutation is somewhat leaky. When cells are cotransfected with pIMRSV-*am* and an amber suppressor tRNA, production of Rep78 is restored to the level observed with pIMRSV. Cotransfection of pIMRSV-*am* with a Rep-expressing plasmid,
10 pRSVrep, provides a high level of Rep78 in *trans*.

Capsid protein expression was analyzed in parallel (Figure 13). Synthesis of the capsid proteins is significantly enhanced following Ad infection of cells
15 transfected with pIM45. This increase is not observed with pIMRSV (lanes 5 and 6) but does occur with the pIMRSV-*am* mutant. The pIMRSV phenotype is restored when pIMRSV-*am* is cotransfected with the suppressor tRNA or when Rep protein is supplied in *trans* by cotransfection with pRSVrep.

20 Northern analysis was performed on samples from the same experiment to examine this phenomenon at the RNA level (Figure 14). With pIM45, a relatively low level of p5 transcript (4.2 kb) was observed, but the p19 transcript
25 (3.6 kb) was more prevalent and showed an increase with Ad infection. The spliced transcripts from p5 and p19 (3.9 and 3.3 kb, respectively) were not detected. Replacement of p5 by the RSV LTR resulted in an increase in the 4.2 kb transcript which contributes to the higher levels of Rep78
30 produced by this construct. Interestingly, the amount of this transcript was augmented further upon introduction of the amber mutation. When synthesis of Rep is restored by cotransfection of a suppressor tRNA (lane 5) or supplied in
35 *trans* by cotransfection with pRSVrep (lane 6) synthesis of the 4.2 kb transcript is again reduced. These results suggest that transcription from the RSV LTR is down-regulated by Rep.

With pIM45, Ad infection results in a significant increase in both the full-length (2.6 kb) and spliced (2.3 kb) p40 mRNA, mirroring the increase in capsid protein synthesis. The prevalence of p40 transcripts over those derived from p5 and p19 is similar to that observed by Labow et al. (1986) from wt AAV in adenovirus-infected KB cells. In general, the ratios of the two p40 mRNAs are shifted in favor of the spliced 2.3 kb transcript with Ad infection. In contrast to pIM45, no increase in the levels of either the p19 or p40 transcripts is observed with the pIMRSV construct upon Ad infection. Notably, with pIMRSV-am, the increase in capsid protein synthesis observed with Ad infection in the Western analysis (Figure 13, lane 4) is not reflected by an increase in the level of cap mRNA. The level of capsid mRNA observed with pIMRSV-am is similar to that of the parental plasmid, pIMRSV. The same lower level is observed in the cotransfection of pIMRSV-am and the suppressor tRNA and of pIMRSV-am with the Rep-expressing plasmid, pRSVrep. The Northern analysis suggests that the reduced level of capsid protein synthesis observed with pIMRSV might be explained at the RNA level by a failure to activate transcription from p40 in response to Ad infection. Furthermore, these results suggest that the increase in capsid protein production following Ad infection with the pIMRSV-am mutant is a post-transcriptional effect and that the mutation has relieved a Rep-mediated translational inhibitory effect on capsid synthesis.

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F. rAAV PACKAGING : COMPARISON OF RSVREP AND CMVCAP
HELPER PLASMIDS.

The series of helper plasmids were compared with respect to their ability to produce rAAV. Each was transfected into Ad-infected 293 cells in conjunction with the pTRLacZ vector plasmid and the yield of rAAV in the

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crude lysate (Table 1) was determined by the titer assay. Increasing capsid protein expression by placing the cap gene under the control of the CMV promoter (p5repΔ-CMVcap) increased rAAV yield by approximately 9-fold. In contrast, replacing p5 with the RSV LTR in order to enhance rep gene expression resulted in a lower rAAV yield. When the RSV LTR was added to a construct containing p40 (pIMRSV or RSV repΔ-p40cap), rAAV titers were decreased by 10-20 fold, while the RSVrepΔ-CMVcap helper packaged rAAV 5-fold less efficiently than the comparable construct containing p5. These results correlate with differences in capsid protein expression observed in the Western analysis. As described above, a dramatic decrease in Cap protein production was observed as a result of overproduction of Rep 78 with the p40-cap constructs, while a more subtle effect was observed on Cap protein expression from the CMV promoter. The results of these experiments comparing the different helper constructs suggest that Cap, but probably not Rep protein synthesis is a limiting factor in the production of rAAV. The Tables provide the results of experiments showing the feasibility of producing recombinant AAV (rAAV) in the absence of adenovirus by providing the required adenovirus early genes (E1A and E1B, E4, E2A, VA) on plasmids.

Table 1: Shows a comparison of rAAV yield from the conventional production scheme (293 standard conditions) in the presence of adenovirus with that obtained in the absence of adenovirus. In the 'absence of adenovirus' cases, several different cell lines were used. Each cell line is a derivative of the 293 cell line (which contains the adenovirus E1 gene) that has been engineered to also contain the adenovirus E4 gene. The VK2-20 cell line was obtained from Frank Graham, while the other ORF 6 lines (2C4 and 3B1) were generated in-house. Since the VK2-20 and ORF6 lines already contain the E1A, E1B and E4 genes, in order to produce rAAV, the E2A and VA genes must be supplied by transfection with the E2A plasmid. E2VA

5' → 3' and E2VA 3' → 5' are two clones of this plasmid (with the insert in opposite orientations). (The cells are also transfected with the rAAV vector plasmid and helper plasmid to allow production of rAAV). The conclusion from this experiment is that it is feasible to produce rAAV in the absence of adenovirus and that this strategy yields as much, if not more, than the conventional method.

Table 2 is a summary of the results of large-scale rAAV production using pIM45 and p5repΔ-CMVcap as helper DNAs. Notably, the yield of rAAV IU/cell increases almost 10-fold when the modified helper is used. This result is also reflected in higher titers (both in IU/ml and particles/ml) of the purified material. Shown in the table are IU/ml determined both in the presence and absence of adenovirus (Adts149). As has been reported by others (Ferrari et al., 1996, Fisher et al., 1996), rAAV titers are approximately 1000-fold higher in the presence of an Ad infection. The particle:IU ratio of these preparations is 20-120 (IU +Ad) or $4 - 7 \times 10^4$ (IU -Ad). The former value is within the range previously reported (Samulski et al., 1989). While the purification procedure results in a persistent yet variable level of Adts149 contamination (from $<10^3$ IU/ml to 10^7 IU/ml;), the stocks are free of contaminating wt AAV (see below).

TABLE 1. Comparison of Helper Plasmids Containing RSV-Rep and CMV-Cap

SAMPLE	TITER (IU/ml)*
pIM45	6.0×10^6
pIMRSV	3.2×10^5
p5rep Δ -CMV cap	5.5×10^7
RSV rep Δ -CMV cap	1.0×10^7
p5rep Δ -p40cap	1.3×10^6
RSV rep Δ -p40cap	1.3×10^5

*Shown is the average of five separate experiments (-lowest, highest values).

293 (standard conditions) - Ad	0
293 (standard conditions) +Ad	4.3×10^6
VK2-20 + E2VA 5'→3' - induction	5.4×10^5
+induction*	7.6×10^5
VK2-20 +E2VA 3'→5' -induction	1.4×10^5
+induction*	7.6×10^5
ORF6 lines +E2VA 5'→3' +induction+	
3B1	3.4×10^7
4B1	4.1×10^7
3C2	2.6×10^6
4A6	8.6×10^6
2C4	1.4×10^8

* lines were induced with dexamethasone (for E4) but not with Zn⁺² (for E2).

+ ORF6 and E2 expression were induced with Zn⁺² and Cd⁺²

TABLE 2. Large Scale Purification of Recombinant AAV(TRIacZ)*

PLASMID	LYSATE		FINAL		
	TITER (IU/ml+Ad)	IU/cell	TITER (IU/ml+Ad)	TITER(IU/ml-Ad)	PARTICLES/ML
pIM45	2.0×10^8	2.9	4.3×10^7	8.0×10^4	7.4×10^9
p5repΔ CMVcap	8.9×10^8	27	5.2×10^8	4.4×10^5	1.9×10^{10}

*shown are the averages of 9 preps for each helper plasmid; these were derived from the transfection of approximately 1×10^9 cells and purified via CsCl gradient centrifugation (see Methods).

H. ANALYSIS OF REPLICATION AND LEVELS OF *wtAAV* CONTAMINATION.

5 Hirt analysis was performed on samples from
small-scale transient transfections such as those described
above to assay replication of the vectors and to assess
levels of *wtAAV* contamination. All of the helper DNAs
supported replication of the TRlacZ vector (Figure 15,
10 panel A); however, in each transfection using a helper
plasmid containing an RSV LTR-*rep* cassette, the vector
appeared to replicate at a diminished level (the
ethidium-bromide stained gel indicated equal amounts of DNA
were loaded in each lane). This result might also help to
15 explain the reduced viral yields obtained with the helpers
containing RSV-*rep*. A low level of replication was
observed when the amber mutant, pIMRSV-*am* was used as a
helper, confirming, as shown in Figure 13, panel A, that a
small amount of full-length Rep protein is synthesized by
20 this mutant. When the same blot was probed with a fragment
from the *wtAAV* genome, no evidence of replicating *wtAAV*
viral DNA was observed (Figure 15, panel C). There was,
however, hybridization to high molecular weight DNA in some
lanes. This could represent cross-hybridization with
25 cellular sequences bearing some homology to the *wtAAV*
probe. However, since no signal appeared in the negative
control (mock) lane, an alternative explanation might be
that the signal is evidence of integration of the helper
DNA into the 293 cell genome. Interestingly, this high
30 molecular weight band appeared only in cells transfected
with helpers carrying the p5-*rep* cassette rather than RSV
LTR-*rep*, suggesting that overexpression of *rep* might
inhibit integration or alternatively, that sequences within
p5 (ie. the Rep-binding site or RRS) might be required in
35 *cis* for integration. Lending support to the integration
hypothesis is the observation that no signal appears in
cells transfected with the mutant pIMRSV-*am*, suggesting

that this phenomenon is dependent upon Rep synthesis. Lysates harvested in parallel from these transfections were used to infect a second plate of 293 cells in the presence of Adts149 and Hirt samples were prepared. If in fact a small amount of contaminating wt AAV were present, then the virus should have been amplified upon reinfection. Southern analysis and hybridization with a wtAAV probe (Figure 15, panel D) again showed no evidence of replicating wtAAV viral DNA. When a duplicate blot was probed with a lacZ fragment (Figure 15, panel B), no replicating vector DNA was observed. This latter result is further evidence of the lack of wt AAV since the presence of wt AAV (i.e., rep gene expression) would have allowed vector replication under these conditions.

AAV promoters in a standard helper plasmid were substituted with stronger heterologous promoters in order to enhance separately the expression of the Rep and Cap proteins required for rAAV packaging. These experiments showed that rAAV yield was improved by approximately 10-fold when cap gene expression was increased, implying that the level of capsid protein is one limiting factor for production of rAAV. In contrast, rep gene expression is probably not a limiting factor since overexpression of rep did not increase rAAV yield. However, it is not possible to make a definitive conclusion on this issue as increases in Rep protein synthesis were always coupled with reductions in capsid protein production. In the case of the plasmid pRSVrepΔ-CMVcap, however, Cap protein production was diminished only slightly relative to that observed with p5repΔ-CMVcap (at most 2-fold, but the level was still higher than attained with pIM45) while Rep78 expression was enhanced significantly (approximately 5-fold). Under these conditions, there was no increase in rAAV yield over p5repΔ-CMVcap; packaging efficiency was in fact slightly reduced. These conclusions conflict with those made based on a previous study (Flotte et al. 1995)

in which use of a construct expressing rep from the HIV LTR (pHIVrep/p40cap) led to a 10-fold increase in rAAV yield compared to a construct in which p5 controlled rep expression (pAAV/Ad).

5 production by the standard factor restricting AAV vector efficiency as the overall protocol is transfection synthesis is limited both by the number of cells taking up DNA as well as the number of DNA molecules present within each cell. In an attempt to increase transfection efficiency, plasmid DNA has been complexed to replication-competent adenovirus modified with polylysine, resulting in an increase in rAAV packaging of 40-240 fold over the standard calcium phosphate method (Mamounas et al., 1995). A number of modifications to the standard rAAV production procedure were made by Chiorini et al. (1995); 10 instead of transfecting 293 cells with calcium phosphate, COS cells were electroporated with reported transfection efficiencies of up to 90%. The helper plasmid used for those studies also contained an SV40 replicon, presumably 15 each transfected cell. By this method, a packaging efficiency of over 10^3 rAAV particles/cell was achieved. Alternatively, packaging cell lines have been constructed in order to avoid the inefficient transfection step. When 20 vector DNA was introduced into a stable cell line, a five-fold improvement in rAAV yield over cotransfection was reported, resulting in 10^4 particles/cell (Flotte et al., 1995). Clark et al. (1995) have constructed a cell line 25 containing both the vector and AAV rep/cap genes which allows production of rAAV by adenovirus infection alone. This system yields 30 IU/cell (approximately 400 particles/cell), a value which is comparable to that achieved with the improved helper plasmid described here. 30 Given the experience of others, it is likely that the packaging protocol employed in these studies can be further 35

improved, either by replicating the helper plasmid within the transfected cell or by using the new helper construct to generate a packaging cell line.

5 The effect observed on Cap protein expression resulting from replacement of p5 with the RSV LTR confirmed the work of others with respect to AAV gene regulation. In addition to their function in replication, the AAV Rep proteins (primarily Rep78/68; Kyostio et al., 1994; Horer et al., 1995) are known to act as transcriptional regulators. In the absence of an adenovirus infection, Rep proteins repress transcription from AAV promoters (Tratschin et al., 1986, Trempe and Carter, 1988, Beaton et al., 1989, Kyostio et al., 1994) while conversely, they activate transcription from these promoters in response to adenovirus infection (Labow et al., 1986, Tratschin et al., 1986). McCartney et al. (1991) have shown that Rep-mediated activation of the p19 and p40 promoters in the presence of adenovirus is dependent in *cis* upon sequences located upstream of both p5 and p19. Consistent with this finding was the lack of induction of p40 transcription upon the RSV LTR infection when p5 was deleted and replaced with the RSV LTR (as in the pIMRSV plasmid). Similarly, there was also no increase in the level of p19 RNA with this construct. The failure to observe induction was due to removal of sequences required in *cis* since it occurred independently of rep gene expression; transcriptional activation of p40 was not restored (Fig. 4) when Rep protein synthesis was prevented by an amber mutation or when Rep protein was supplied by an amber mutation or pIMRSV lacks only 84 bp upstream of p5 (bp 191-275 of the AAV sequence); this deletion is more limited than the one reported by McCartney et al. (1991) (bp 191-320) and thus further defines the location of the putative regulatory region required for Rep activation. The region between bp 191 and 275 is known to contain binding sites for the major late transcription factor (USF; Chang et al., 1989), YY1

(Shi et al., 1991) and Rep (McCarty et al., 1994; Kyostio et al., 1995) as well as the p5 TATA box.

5 Though transcription from the p40 promoter in the
pIMRSV-*am* mutant was not activated by Rep in response to
adenovirus infection, Cap protein synthesis was observed to
increase. This effect may be attributed to the
translational inhibitory activity of Rep. In 293 cells in
10 the absence of an adenovirus infection, Trempe and Carter
(1988) observed that the level of p40 mRNA was reduced
while CAT protein expression increased in the absence of
Rep compared to a *rep* gene-containing vector. In cells
transfected with pIMRSV*am*, synthesis of the capsid proteins
is significantly enhanced with adenovirus infection. This
15 increase, however, occurs without any alteration in the
steady-state level of p40 mRNA, indicating that it is a
translational effect. In comparison, capsid protein
production also increases in cells transfected with pIM45,
but in this case, there is a concomitant increase in the
20 level of both the 2.6 kb and 2.3 kb p40 mRNAs. The
apparent induction in the synthesis of the capsid proteins
with pIMRSV*am* is a *trans* effect of the mutation of the *rep*
gene, as it does not occur in any case where the Rep
proteins are expressed. Because Rep78 is the major Rep
25 protein produced by pIMRSV, it is presumably the primary
mediator of the inhibitory effect, however, a role for
Rep68 cannot be ruled out. These results suggest that
although adenovirus infection is capable of significantly
increasing the efficiency of translation of p40 mRNA (West
30 et al., 1987; Janik et al., 1989), this effect can be
counteracted by the Rep proteins. It is not clear whether
translational inhibition in the presence of adenovirus
occurs as a normal function of Rep or if it is an artifact
of overexpression of the *rep* gene in the pIMRSV construct.
35 Alternatively, inhibition may occur only when the level of
p40 mRNA is low and can be overcome when transcription from
p40 normally increases with adenovirus infection.

Induction of transcription from p40 was prevented in this case by removal of sequences upstream of p5.

These experiments have provided further evidence of the ability of the Rep proteins to act as repressors of expression from heterologous promoters. The Rep proteins are known to down-regulate expression of several heterologous genes (Labow et al., 1987; Antoni et al., 1991; Rittner et al., 1992; Oelze et al., 1994; Horer et al., 1995). In the experiments described here, expression of the cap gene from the CMV IE promoter as well as the rep gene from the RSV LTR were both down-regulated by Rep. Rep has previously been shown reduce the level of expression of the cat gene from the CMV IE promoter (Heilbronn et al., 1990); similar to the results obtained here, this effect was minor (approximately 2-fold). For both the CMV IE and RSV LTR promoters, inhibition occurred at the RNA level, though since steady-state levels of RNA were assayed by the Northern analysis, the effect could be either at the level of transcription or mRNA stability. Down-regulation at the RNA level has also been demonstrated in the case of the HIV LTR and HPV18 URR promoters and has been attributed primarily to Rep 78/68 (Antoni et al., 1991; Horer et al., 1995).

The following examples are intended to illustrate the invention without limiting the scope thereof.

EXAMPLE 1: RECOMBINANT AAV PRODUCTION USING A NON-REPLICATING HELPER PLASMID CONTAINING AAV HELPER GENES

An experiment was performed to determine if a helper plasmid providing the AAV rep and cap genes on a nonreplicating plasmid and using heterologous promoters for gene expression would increase AAV vector titer above that derived from controls in which the rep and cap genes are

expressed from their own promoters. The helper plasmids tested are shown in FIGURE 7.

5 pIM45 contains the AAV *rep* and *cap* genes under the control of the native AAV promoters, and the AAV polyA site at the 3' end (McCarty, D.M. et al., *J.Virol.* 65:2936-2945, 1991).

10 pRSVrep-p40cap and pRSVrep-CMVcap were constructed by deleting the *cap* region from pIM45 using PCR to delete nucleotides 2287-4049 in the AAV genome, resulting in the generation of pRep*30. A PCR fragment isolated from pRep*30, nucleotides 275-4464, containing *NheI* and *NotI* ends, was cloned in between the *NheI* and *NotI* sites in pRep9 (Invitrogen, San Diego, CA) to make pRSVrep. 15 An *XbaI* (filled in)-*SfiI* fragment from pRSVrep was cloned into *SmaI* and *SfiI* digested pIM45 to make pRSVrep-p40cap. This same *XbaI* (filled in)-*SfiI* fragment from pRSVrep was cloned into *SmaI* and *SfiI* digested pIM-CMVcap (see below) 20 to make pRSVrep-CMVcap.

pIM-CMVcap was constructed by introducing 3 point mutations at positions 1823 (T:C), 1826 (A:C) and 1832 (G:A) within the AAV genome to inactivate the p40 promoter. 25 pCMVcapB was generated by inserting a PCR fragment containing nucleotides 1850-4460 from the AAV genome and *BamHI* ends into the *BamHI* site of pCMVB (Clontech, Palo Alto, CA). An *SphI* fragment containing the CMV promoter was isolated from pCMVcapB and was inserted into the *SphI* site at position 930 within pIM45 to make pIM-CMVcap. 30

The helper plasmids were transfected into 293 cells at a ratio of 10:1 helper/vector (16.5 µg/total DNA). The AAV vector plasmid used was pTR-lacZ, which was 35 developed by Dr. Nicholas Muzyczka, University of Florida. The plasmid is shown in FIGURE 8. Isolation and

purification of the AAV vector was performed as described in Section 4, *supra*.

The AAV yield was titered by coinfecting 293
5 cells with helper adenovirus and the AAV. This reduced the
infection time, therefore increasing the sensitivity of the
assay. For the titer, 293 cells were plated in 96-well
plates, at 5×10^5 cells/ml. (100 μ l/well) in DMEM/10% FBS
(with penicillin/streptomycin and glutamine) and allowed to
10 grow for one day at 37°C. The cells were then coinfectd
with Adts149 virus using an MOI of 20, and with the AAV at
dilutions of 1:100, 1:200 and 1:400 etc. of the viral
preparatory stock. The different dilutions were used in
order to ascertain the titer.

15 After the infection was allowed to progress for
two days at 39°C, the medium was aspirated, the cells were
incubated with 3.7% formaldehyde for 5 minutes and washed
with Phosphate Buffer Saline (PBS). The cells were then
20 stained with X-Gal (5-Bromo-4-Chloro-3-Indolyl- β -D-
galactopyranoside) at 37°C for 1-24 hours and screened for
the presence of blue coloration in cells in order to detect
the expression of the *lacZ* gene contained in the AAV
vector. A conversion using the Titer Analysis program,
25 which is based on determining the endpoint dilution, was
used for determination of the titer in IU/ml.

A Western blot analysis was performed to
determine the levels of rep and cap protein expression
30 using the various helper plasmids (and specific clones)
shown in FIGURE 7. FIGURE 9 shows a Western blot analysis
of rep protein expression, while FIGURE 10 shows a Western
blot analysis of cap protein expression. Standard
techniques were used in the Western blot analysis (Current
35 *Protocols in Molecular Biology*, Ausubel, F. et al., eds.,
Wiley and Sons, New York, 1995).

The titer data is shown in TABLE 3. The titer of vector stock is given in IU/ml. The experiment demonstrates that increased levels of cap expression in particular, as evidenced by the Western blot, leads to increased production of the AAV vector, pTRLacZ, as evidenced by the titers shown.

TABLE 3

10

COMPARISON OF HELPER PLASMIDS CONTAINING RSV-REP AND CMV-CAP:LYSATE TITERS	
SAMPLE	TITER (IU/ML)
mock	0
pIM45	1.7×10^6
pIM-CMVcap	3.2×10^7
pRSVrep-p40cap clones	
1-8	2.4×10^6
2-1	2.2×10^5
3-14	3.6×10^5
8-1	1.5×10^5
pRSVrep-CMVcap clones	
1-10	1.4×10^7
2-4	4.9×10^6
3-12	1.9×10^7
8-14	1.2×10^7

EXAMPLE 2: CELL LINES, VIRUSES AND PLASMID DNA.

5 The 293 cell line, an adenovirus 5-transformed
human embryonic kidney cell line (Graham et al., 1977) was
propagated in Dulbecco's modified Eagle's medium-high
glucose (DME; Irvine Scientific) supplemented with 10%
fetal bovine serum (FBS; Irvine Scientific, Santa Ana, CA),
20 mM glutamine, 100 units/ml penicillin and 100 µg/ml
10 streptomycin (Gibco-BRL, Gaithersburg, MD) at 37°C and 5%
CO₂. The adenovirus type 5 mutant, ts149 (Ad5ts149;
Ensinger and Ginsberg, 1972) used as a helper virus in
these studies has reduced ability to replicate viral DNA at
the nonpermissive temperature (39°C) due to a
15 temperature-sensitive mutation in the DNA polymerase
encoded by adenovirus early region 2 (Stillman et al.,
1982). Ad5ts149 was grown in 293 cells at the permissive
temperature (33°C) and purified by CsCl gradient
centrifugation.

20 Plasmid DNA encoding the recombinant AAV vector,
pTRLacZ, as well as the helper plasmid, pIM45 (McCarty et
al., 1991), were generously provided by N. Muzyczka (Univ.
of Florida). pTRLacZ consists of the *E. coli* LacZ gene
25 (cytoplasmic) under the transcriptional control of the CMV
IE promoter, inserted between the terminal repeats of AAV.
The plasmid encoding the amber suppressor tRNA, pSVtsSu⁺
(amber) (Capone et al., 1985) was obtained from U. L.
RajBhandary (MIT). pNTC3, an AAV genomic clone containing
30 an amber mutation within the rep coding region (Chejanovsky
and Carter, 1989) was kindly provided by R. Owens (NIH).

A. PLASMID CONSTRUCTION.

35 Using pIM45 as the starting plasmid, the
endogenous AAV promoters, p5 and p40 were replaced with the

Rous Sarcoma Virus long terminal repeat (RSV LTR) and CMV IE promoter, respectively. All manipulations were carried out following standard cloning procedures (Sambrook et al., 1989). All restriction and DNA-modifying enzymes were
5 obtained from New England Biolabs (Beverly, MA) and used according to the manufacturer's specifications. Plasmid DNAs were purified using kits obtained from Qiagen (Chatsworth, CA).

10 The CMV IE-cap cassette was constructed by first amplifying a DNA fragment consisting of AAV genomic sequences between bp 1852 and 4440 (encoding the capsid proteins and including the AAV mRNA polyadenylation site) via PCR (Saiki et al, 1988) using Vent polymerase (New
15 England Biolabs, Beverly, MA). This fragment was inserted between the *Bam*HI sites of pCMV β (Clontech, Palo Alto, CA) to generate the plasmid pCMVcap.

To derive a minimal Rep-encoding sequence, *rep*
20 gene sequences between the *Bam*HI site (bp 1045) and *Apa*I site (bp 2283) were PCR-amplified and inserted within the pIM45 plasmid digested with *Bam*HI and *Apa*I. The result was a deletion between bp 2283 (just downstream of the Rep termination codon) and the *Apa*I site at bp 4049. This
25 plasmid, pIMrep Δ , was used to generate a construct in which Rep78/68 are expressed from the RSV LTR. A 2.4 kb *rep* gene fragment extending from bp 276 (just upstream of the Rep78/68 mRNA initiation codon) to bp 4459 was
PCR-amplified from pIMrep Δ and inserted between the *Nhe*I
30 and *Not*I sites of the pRep9 expression vector (Invitrogen, San Diego, CA) to create pRSVrep.

Because the Rep and Cap protein coding sequences overlap in the region of the AAV intron, there is 431 bp in
35 common between the *rep* and *cap* gene cassettes (between bp 1852 and 2283) of pIMrep Δ and pCMVcap. Prior to insertion of the CMV IE-cap fragment into pIMrep Δ to create

p5repΔ-CMVcap, p40 sequences within pIMrepΔ were mutated to inactivate the promoter. This was done to prevent the generation of wild-type AAV as a consequence of recombination between the shared sequences. Mutagenesis was carried out by overlap extension PCR (Higuchi et al., 1988). pIMrepΔ was used as a template for the first PCR using flanking primer-1 (5'-GGATTACCTCGGAGAAGCAGTGGATCC-3'; bp 1024-1050 of the AAV genome) and mutagenic primer-1 (5'-GTTTGGGTTCACTGATGCTGCGTCACTG-3'; AAV bp 1821-1841; mutated nucleotides are underlined). The result is the introduction of three base pair mutations in the region of the p40 TATA box : from TATAAGTGAG to CATCAGTGAA. The G to A change ablates a *BanII* site to enable screening by restriction analysis. pIMrepΔ was also used as a template for the second PCR using flanking primer-2 (5'-GTGTGGAATCTTTGCCCAGATGGGCCCGTTT-GAGCTTC-3'; AAV bp 2260-2283, 4049-4066) and mutagenic primer-2 (5'-CAGTGACGCAGACATCAGTGAACCCAAACG-3'; AAV bp 1821-1841). After gel purification of the above PCR products, a third PCR was performed by annealing the two earlier products and carrying out a final amplification step using only the flanking primers, thereby generating a 1285 bp DNA fragment. This fragment was digested with *BamHI* and *ApaI* and cloned into the corresponding sites of the pIMrepΔ backbone. The resulting plasmid was pIMrepΔ/p40Δ. The helper plasmid p5repΔ-CMVcap was constructed by inserting a *SphI* fragment from pCMVcap containing the CMV IE promoter and *cap* gene cassette into the unique *SphI* site of pIMrepΔ/p40Δ. Similarly, in order to construct p5repΔ-p40cap, a PCR fragment with *SphI* ends extending from AAV bp 1715 to 4461 was generated from pIM45 and cloned into the *SphI* site of pIMrepΔ/p40Δ.

The p5 promoter regions in the plasmids pIM45, p5repΔ-CMVcap and p5repΔ-p40cap were replaced with the RSV LTR promoter by first cleaving pRSVrep with *XbaI*. The *XbaI* site was made blunt with DNA PolymeraseI-Klenow fragment

and the DNA was restricted with *SfiI* to release a fragment containing the RSV promoter and the 5' end of the rep gene. This fragment was then cloned between the *SmaI* and *SfiI* sites of the parental plasmid.

5

To introduce an amber mutation into pIMRSV, a *SfiI*-*BamHI* fragment containing the mutation (at bp 1033 of the AAV genome) was isolated from the plasmid pNTC3 (Chejanovsky and Carter, 1989) and cloned into the corresponding sites of pIMRSV.

10

EXAMPLE 3: TRANSIENT TRANSFECTIONS AND ANALYSIS OF rAAV REPLICATION AND PACKAGING.

15

For small-scale experiments, 293 cells were seeded at a density of 1×10^6 cells per 6 cm dish 48 hours before transfection. The cells were infected with Ad5ts149 in DME-10% FBS at a multiplicity of infection (MOI) of 20 for 1 hour at 37°C prior to transfection. Transfection procedures were carried out using the calcium phosphate Profection kit (Promega, Madison, WI) according to manufacturer's instructions. In general, for rAAV packaging, each dish received a mix of 1.5 µg vector DNA (i.e. pTRLacZ) and 15 µg of helper DNA. Following incubation at 37°C for 5 hours, the infection/transfection was terminated by replacing the media with fresh DME-10% FBS; the dishes were then transferred to 39°C (the non-permissive temperature for Ad5ts149).

20

25

30

For analysis of rAAV packaging, cells were harvested at 48 hours post-transfection by low-speed centrifugation in a clinical centrifuge. The pellet from each dish was resuspended in 100 µl phosphate-buffered saline (PBS) and freeze-thawed four times to liberate the rAAV. Adenovirus was heat-inactivated by incubating the lysate at 56°C for 30 minutes. The lysate was subjected to

35

a second low-speed spin to pellet cellular debris and the supernatant was collected. The rAAV titer was determined on 293 cells (+/- coinfection with Ad5ts149; MOI=20) by endpoint dilution. Following staining of the cells with
5 X-gal (5-bromo-4-chloro-3-indoyl- β -D-galacto-pyranoside) for 20-24 hours, titers were calculated using a computer program based on Karber's method (Lynn, 1992).

Replication of vector DNA in transfected cells
10 was assayed by isolating extrachromosomal DNA 48 hours post-transfection according to the Hirt fractionation method (Hirt, 1967). DNAs were restricted with *DpnI* (to digest input DNA) prior to agarose gel electrophoresis and Southern analysis. The *lacZ* and wild-type AAV probes used
15 were both 50-mer oligonucleotides, 5'-ACTGCTGCCAGGCGCTG ATGTGCCCCGGCTTCTGACCATGCGGTTCGCGTTC-3' and 5'-TCGGAGGAAGC AAGGTGCGCGTGGACCAGAAATGCAAGTCCTCGGCCAG-3' (AAV nucleotides 1501-1550), respectively. These were labelled with [γ -³²P]
ATP using T4 polynucleotide kinase according to standard
20 procedures (Sambrook et al., 1989). The filter was hybridized and washed as described below for the Northern blot analysis, except that the prehybridization, hybridization and final wash step were at 60°C.

25 A. PROTEIN EXTRACTION AND IMMUNOBLOTTING.

For analysis of Rep and Cap protein expression from the various helper plasmids, 293 cells were first transfected as described above. Nuclear fractions were
30 prepared 48 hours post-transfection according to the procedure described by Mendelson et al. (1986). Sample volumes were normalized according to DNA content (by optical density at 260 nm), mixed with 15-20 μ l of sample buffer (500 mM Tris-HCl, pH 6.8, 10% sodium dodecyl sulfate
35 (SDS), 20 mM EDTA, 10% β -mercaptoethanol, 10% glycerol, and

0.2% bromophenol blue), and boiled for 5 minutes prior to loading.

Following electrophoresis in 10%

5 polyacrylamide/0.1% SDS gels, proteins were transferred from the gel to Hybond polyvinylidene difluoride (Amersham, Arlington Heights, IL) membranes. Prior to staining, the filters were blocked for 1 hour at room temperature in 5% milk powder dissolved in TBST (10 mM Tris HCl, pH 8.0, 150
10 mM NaCl and 0.05% Tween 20). The primary antibodies used for Rep and Cap Westerns were both mouse monoclonals (American Research Products, Belmont, MA): anti-AAV Rep protein, 303.9 (used at a dilution of 1:10 in TBST) and anti-VP1, VP-2 and VP-3 of AAV, B1 (used at a dilution of
15 1:5 in TBST), respectively. These were incubated on the filter for 2 hours at room temperature with vigorous shaking. Following a wash step in TBST (3 x 15 minutes), the filter was incubated in the secondary antibody, goat anti-mouse IgG (Fab-specific) peroxidase conjugate (Sigma,
20 St. Louis, MO), for 1 hour at room temperature. The filter was then washed as before and developed using the ECL kit (Amersham).

B. ISOLATION OF RNA AND NORTHERN ANALYSIS.

25

Total RNA was isolated from transfected 293 cells using RNazol B (Tel-Test, Inc., Friendswood, TX) according to the manufacturer's instructions. Prior to
electrophoresis, 10 µg of each RNA was combined with
30 denaturation cocktail (50% DMSO, 10% formaldehyde, 20 mM MOPS (morpholinepropanesulfonic acid), pH 7.0, 10 mM sodium acetate, 1 mM EDTA) and loading dyes (5% glycerol, 0.1 mM EDTA, 0.04% bromophenol blue, 0.04% xylene cyanol) and
heated at 65°C for 15 minutes. Electrophoresis was through
35 a 1% agarose/ 0.65% formaldehyde gel assembled and run in MOPS running buffer (20 mM MOPS, pH 7.0, 10 mM sodium acetate, 1 mM EDTA). Transfer to GeneScreen nylon membrane

(NEN-DuPont, Boston, MA) was carried out by capillary action overnight in 10 x SSC (1.5 M NaCl, 0.15 M sodium citrate; Sambrook et al., 1989).

5 The filters were prehybridized for 4-5 hours at 65°C and then hybridized with probe overnight at 65°C in hybridization buffer (5 x SSC, 0.5% SDS, 5 x Denhardt's solution (Sambrook et al., 1989), 100 µg/ml denatured salmon sperm DNA). The probe was a 1.6 kb HincII fragment
10 of pIM45 (AAV bp 2397 to 3987) labelled with [α -³²P]dATP (specific activity, 3,000 Ci/mmol; NEN-DuPont, Boston, MA) using a random primer labelling kit (Stratagene, La Jolla, CA). The filter was washed for 5 minutes at room temperature in 2 x SSC, 0.5% SDS, 15 minutes at room
15 temperature in 2 x SSC, 0.1 % SDS, and then for 2 hours in 0.1 x SSC, 0.5 % SDS at 65°C and exposed to film.

C. LARGE-SCALE TRANSFECTION AND rAAV PURIFICATION.

20 Prior to transfection of 293 cells for large-scale growth of rAAV the cells were seeded in roller bottles such that they would reach 60-80% confluence on the day of transfection (final density was approximately 1 x 10⁸ cells/bottle). Transfection was carried out in OptiMem
25 media (Gibco-BRL Life Technologies, Gaithersburg, MD) using lipid #53:DOPE (Lee et al., 1996), 22 µg of vector DNA and 218 µg of helper DNA per bottle. Cells were infected with Ad5ts149 at a MOI of 20 at the time of transfection and incubated at 39°C for 48 hours prior to harvest.

30 At the time of harvest, cells were dislodged from the bottles by gentle shaking. The cells were pelleted by centrifugation in a Sorvall RC-3B swinging bucket rotor (2500 rpm, 4°C, 15 minutes) and frozen. For purification
35 of rAAV, the cells were resuspended in PBS containing 2 mM MgCl₂, 0.7 mM CaCl₂, 10% glycerol, and 0.1% Tween.

Benzonase® (Nycomed Pharma A/S, Copenhagen, Denmark) was added (10 µl/1 x 10⁸ cells) and the suspension was incubated with shaking for 1 hour at room temperature. Trypsin (Gibco-BRL Life Technologies) was added to a final concentration of 0.25% and the suspension was incubated again with shaking for 1 hour at room temperature. The cell debris was collected by centrifugation (3000 rpm, 15 minutes, 4°C in Sorvall RC-3B) and the lysate was filtered through a 0.45 µm filter.

10

The lysate was then subjected to centrifugation through a CsCl step gradient (4 hours, 26K rpm, 4°C, SW28 rotor) in which the top and bottom layers were 1.37 g/ml and 1.5 g/ml CsCl, respectively. The top layer was collected (between the CsCl interface and the Ad5ts149 band), adjusted to 1.41 g/ml CsCl, and centrifuged through a 1.41g/ml CsCl equilibrium gradient (16-20 hours, 4°C, 35,000 rpm, NVT.65 rotor). Fractions were collected and assayed on a refractometer; fractions with a density of 1.36-1.41 were pooled and dialyzed against PBS/1% sucrose for 6 hours at 4°C. Sucrose was added to a final concentration of 5% and the purified virus was stored in aliquots at -80°C.

25

D. CHARACTERIZATION OF PURIFIED rAAV STOCKS.

The purified rAAV stock was titered for rAAV in the presence and absence of Ad5ts149 (MOI=20) by endpoint dilution as described above. The titer of contaminating Ad5ts149 was determined in a similar manner except staining was for hexon using anti-adenovirus (hexon) /FITC conjugate (Chemicon, Temecula, CA). The level of contaminating wild-type AAV was assayed using the infectious center assay as described (Einerhand et al., 1995).

35

AAV particle titer was quantitated using a procedure modified from Samulski et al. (1989). The purified rAAV sample was first treated with proteinase K in 0.1% SDS at 37°C for 3 hours. Appropriate dilutions as well as standard curve DNAs (for TRlacZ virus, pTRLacZ DNA was used as a standard) were treated with denaturation solution (0.5 M NaOH, 1.5 M NaCl) for 10 minutes at room temperature and a 1 ml volume was applied to a GeneScreen Plus (Amersham) membrane using a slot blot apparatus (Schleicher and Schuell, Keene, NH). After loading, the slot was washed with 300 µl of 0.5 M ammonium acetate, pH 5.2. The filter was dried and hybridized as described above. The probe (a PvuII fragment of pTRLacZ) was labelled using the Prime-It Fluor labelling kit (Stratagene, La Jolla, CA). Following a series of washes as described above (except the final wash at 65°C was for 10 minutes), the filter was developed with the Illuminator detection kit (Stratagene). Particle concentrations were estimated by comparing the sample signal with that of the standard curve.

EXAMPLE 4: GENERATION OF CELL LINES

293-MT-DBP: Plasmid Construction. The parental plasmid pREP-7 (Invitrogen, San Diego, CA), contains the EBV origin of replication and the EBNA-gene for plasmid extrachromosomal maintenance and the hygromycin resistance gene for DNA selection. To construct pREP/MT/DBP, the RSV promoter of pREP-7 was replaced with the metallothionein (MT) promoter, which is induced by heavy metals. The E2A gene encoding DNA binding protein (DBP) was cloned downstream of the MT promoter.

Generation of 293/MT/DBP Cell Line. 293 cells were transfected with pREP/MT/DBP clone 0.5 via calcium phosphate transfection. Transfected cells were selected for 6 weeks with hygromycin; selected cells were clonally

expanded and judged on the basis of DBP expression (via immunofluorescence) after induction and ability to complement E1-/E2A vectors.

5 3B1 and 2C4 Cell Lines: The parental plasmid contains an expression cassette for adenovirus E4 6 and 6/7 open reading frames (ORFs). The promoter used to drive expression is a mutant human metallothionein promoter which has a low level basal activity. (Makarov et al., *Nuc. Acids*
10 *Res.* 22(8):1504-1505 (1994)).

Both 3B1 and 2C4 cell lines were derived from 293 cells which are human embryonic kidney cells that have been transformed with the E1 region of Adenovirus Type 2. Both
15 3B1 and 2C4 have the ability to complement recombinant adenovirus vectors that are deleted for E1 and E4. The cell lines contain a mutant human metallothionein promoter and SV40 splice and polyadenylation signals to drive expression of adenovirus type 2 E4 open reading frames 6
20 and 6/7 (adenovirus nucleotides 34082-32913). For complementation of E4 functions, the expression of ORFs 6 and 6/7 can be induced by the addition of 100 μ M Zn²⁺, 2 μ M Cd²⁺. Briefly, 293 cells were transfected with the parental plasmid. The cells were cotransfected with
25 pSV2Neo so that individual clones could be selected with G418.

EXAMPLE 5: TRANSGENIC MICE WITH AAVS1 INTEGRATION LOCUS.

30 Characterization of a preferred site on human chromosome 19 for integration of adeno-associated virus DNA by non-homologous recombination. 700-800 CD-1 mice were injected with a purified DNA fragment (0.7 kb *EcoRI-SacI* fragment of AAVS1; Kotin et al., *EMBO J.*
35 11:5071-5078). 550-600 eggs survived and were cleaved. 19 mice were implanted with injected eggs and 148 pups were born.

Chromosomal DNA was isolated from mouse tails and was screened by Southern analysis. Six positive mice were found (#66, 73, 85, 93, 123, 147) (Table 4).

5

Table 4. Screening of F₀ mice

Mouse #	Copy #	Sex	Restriction Enzyme(s)
66	<5	male	AvaI
73	<5	female	AvaI, BamHI
85	<5	male	AvaI, BamHI
93	<5	female	AvaI
123	10-20	female	AvaI
147	<5	female	AvaI
Note: Copy # estimated due to probable mosaic nature of mice			

10

Positive F₀ were mated with wt CD-1 mice and produced seventy-two F₁ pups. Chromosomal DNA from F₁ progeny was isolated and was screened using PCR (250 bp fragment produced by positive samples in the presence of DMSO) (Table 5). Based on the results of the screening, a

15

total of 43 mice were kept (Table 6).

Table 5. Screening of F1 progeny

F ₀	Sex	# of pups	Positive by PCR			Negatives Kept	
			Total	Sex:#	(%)	Total	Sex:#
#66	M	27	9	F:6 M:3	(33)	2	F:1 M:1
#73	F	14	11	F:6 M:5	(79)	2	F:1 M:1
#85	M	5	4	F:0 M:4	(80)	1	M:1
#123	F	12	5	F:1 M:4	(29)	3	F:1 M:2
#147	F	14	4	F:3 M:1	(29)	2	F:1 M:1
TOTAL	6	72	33			10	
			TOTAL # MICE KEPT = 43 (pos.+ and neg.-)				

5

TABLE 6

AAV-1-66 M	AAV-1-73 F	AAV-1-85 M	AAV-1-123 F	AAV-1-147 F
Offspring # (Sex) +/-	Offspring# (Sex) +/-	Offspring # (Sex) +/-	Offspring # (Sex) +/-	Offspring # (Sex) +/-
149 (F) -	176 (F) +	190 (M) +	195 (M) -	205 (M) -
154 (F) +	177 (F) +	191 (M) +	196 (M) +	208 (M) +
155 (F) +	178 (F) +	192 (M) -	197 (M) -	211 (F) +
157 (F) +	179 (F) +	193 (M) +	200 (M) +	212 (F) +
159 (M) -	180 (F) -	194 (M) +	201 (M) +	213 (F) +
164 (M) +	181 (F) +		202 (F) +	214 (F) -
165 (M) +	182 (F) +		203 (F) -	
170 (M) +	183 (M) -			
171 (F) +	184 (M) +			
172 (F) +	186 (M) +			
174 (F) +	187 (M) +			
	188 (M) +			
	189 (M) +			

A. ESTABLISHMENT OF PRIMARY CULTURES FROM F1 TRANSGENICS.

5

Briefly, tail samples were chopped and trypsinized as with previous samples. Free cells and cell chunks were placed 6-well dishes (one well/tail) in 10% calf serum in DMEM. Analysis of cell 4 days after indicated several attached cells had a fibroblast-like morphology. At that point the cell chunks were removed and replaced with fresh media. Cells from the primary F1 cultures are infected with AAV (10 MOI) (AAV titer 9 x

10

10E9/ml = 9×10^6 /ul) (1×10^6 cell/plate). 48 hours post-infection cells are harvested and subjected to Hirt analysis.

5 Table 7. Evaluation of Chromosomal Integration

MOUSE # (SEX)	PARENT	INTEGRATION LOCUS	
		Positive +	Negative -
164 (M)	AAV-1-66	+	
165 (M)		+	
159 (M)			-
184 (M)	AAV-1-73	+	
186 (M)		+	
190 (M)	AAV-1-85	+	
191 (M)		+	
196 (M)	AAV-1-123	+	
199 (M)		+	
195 (M)		-	
208 (M)	AAV-1-147	+	
211 (F)		+	

10 Although the invention has been described with reference to the disclosed embodiments, it should be understood that various modifications can be made without

departing from the spirit of the invention. Accordingly,
the invention is limited only by the following claims.

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